

FEATURES		Location/Qualifiers	
source	1. .475		
/organism="Homo sapiens"			
/note="Organ: colon; Vector: pBluescript SK-; Site_1:			
EcoRI; Site_2: XhoI"			
/db_xref="ATCC (inhost):108677"			
/db_xref="taxon:9606"			
/clone_lib="colon carcinoma (HCC) cell line"			
/tissue_type="colon"			
/cell_type="KM12C"			
/cell_line="KM12C(HCC)-parental human colon			
carcinoma;Dukes B2"			
mrna	<1. >475		
BASE COUNT	108 a 107 g 135 t	6 others	
ORIGIN			
Query Match 54.9%; Score 444.4; DB 17; Length 475;			
Best local Similarity 98.1%; Pred. No. 1.4e-114;			
Matches 467; Conservative 0; Mismatches 7; Indels 2; Gaps			
QY	189	GGCGGGCGCGTGGCGGTCTGCTCGGAGGCGGTGACGGCCAGACTCGTTGG	248
DB	1	GGCGGGCGCGTGGCGGTCTGCTCGGAGGCGGTGACGG-CAGACTCGTTGG	59
QY	249	TGTCCTGTGTTGCTCTCAGTCACTACAGAACCTCGGGGGCTGTTCACCTCGCGGG	308
DB	60	TGTCCTGTGTTGCTCTCAGTCACTACAGAACCTCGGGGGCTGTTCACCTCGCGGG	119
QY	309	GGCGGAGGAGTCCTTAAGTGGGAGACCTCAAAGTGGGACAATATTTGTAAGATCC	368
DB	120	GGCGGAGGAGTCGCTTAAGTGGGAGACCTCAAAGTGGGACAATATTTTAAAGATCC	179
QY	369	AAAAATAATGAGCTACGCAAGAACCTTAAGTACAACTACACAGCTCATGTTTC	428
DB	180	AAAAATAATGAGCTACGCAAGAACCTTAAGTACAACTACACAGCTCATGTTTC	239
QY	429	CTGTTTCCAGCACCACAACTAATCTTAAGGATCCAGTGGCAATGAACACATTTTAC	488
DB	240	CTGTTTCCAGCACCACAACTAATCTTAAGGATCCAGTGGCAATGAACACATTTTAC	299
QY	489	TGGGACGAATTTGGTTTTTCAAGCCCATATCTTCCCAATATTAATGGCTATTCCTA	548
DB	300	TGGGACGAATTTGGTTTTTCAAGCCCATATCTTCCCAATATTAATGGCTATTCCTA	359
QY	549	CAAAATGGCAGTCGCATTTCTTTTCTTGGATGTTGGGAGCAGATCGATTTTACCT	608
DB	360	CAAAATGGCAGTCGCATTTCTTTTCTTGGATGTTGGGAGCAGATCGATTTTACCT	419
QY	609	TGGATACCTCGCTTTGGGTTGTTAAAG-TTTTCAGCTAGAGGTTTTCGGAATT	663
DB	420	TGGATACCTCGCTTTGGGTTGTTAAAGTTTTCAGCTAGAGGTTTTCGGAATT	475
RESULT	4		
AI299154			
LOCUS	GM96f11.x1 NCI_CGAP_Lu5 Homo sapiens cDNA clone IMAGE:1896621	3'	03-DEC-1998
DEFINITION	mrna sequence.		
ACCESSION	AI299154		
NID	9395808		
KEYWORDS	EST.		
SOURCE	human.		
ORGANISM	Homo sapiens		
REFERENCE	1 (bases 1 to 439)	Eukaryota; Metazoa; Chordata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.	